Figure 1. Amino acid sequence alignment in human IgG isotypes and their variants.

	Amino Acid Position								
Human IgG Isotype	228234	235	236	237330	331				
G1	ProLeu	Leu	Gly	GlyAla	Pro				
G2	ProVal	Ala		GlyAla	Pro				
G4	SerPhe	Leu	Gly	GlySer	Ser				
G1 variant	Pro Val	Ala	Gly	GlyAla	Ser				
G2 variant	ProVal	Ala		GlyAla	Ser				
G4 variant	<i>Pro</i> Phe	Ala	Gly	GlySer	Ser				

5	ID number	Corresponding Row in this Figure 1
	SEQ ID NO:26	G1
	SEQ ID NO:27	G2
10	SEQ ID NO:28	G4
	SEQ ID NO:22	G1 variant
15	SEQ ID NO:18	G2 variant
	SEQ ID NO:20	G4 variant

acid sequences of hG-CSF-L-vFcy2 amino deduced

9 120 180 240 300 360 420 480 540 009 099 cag Q gag E CCC tac Y aca T cag Q အီဝီင လ gga G tac Y cct P gac D gcc A atc I tgc C ctc ctg L ttc F CCC P tcg S ggc G aag K agc S ttc F ttg L gаа Е gct A gtg V ggc G ctg L а**g**g R tga C agc S ctt L acc T gaa E tct s gag E gga G atg M gcc A ctg L ggc G CCC P ctg L ggt G ctg L gct A ttg L gcc A CCC P gag E tac Y cag Q agc S cat H ggt G tgg W gag E agc S cag Q tta L acc T ctc tgg W ccg P ggc G cag Q tgc C CCC CCC P atg M gcc caa Q atc I ggt G gtg V ctg L gcc A acc T aca T a a K tgt C atc I agc S tcc s acc H cat H tcc S gga G gcc A ggc G atc I ggt G ctc acc T tgg W ctg L ttg L tcc S ctg L 999 G gcc A cct P ctc aag K ctg L tgc C gcc A cag Q CCC P ttc F tct S ggc acc T gca A gag E gаа Е ttt F gtt V cag Q အီဝိုင လ cag Q CaC H gca A ctg L gac D CCC P cta L gcc A atg M -30 gcc A cac H -10 cag gga G ctg L gcc A cag Q gtc V ctt L ctc tgg W cac H 170 ctg L ctg L ctg L ctg L gca A ctg L gac D gcc gga G cgc R ctc ctg L tcc S gtg V cct ctg L ggc G gcc gca A agc S 999 G ctg L ctg L gat D cag Q gcc A cag Q gtt V cgg R cag Q gcc ggc G gag E agc S cag Q ctg L atg M aga R ng D R

-4												
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DNA and deduced amino acid sequences of hG-CSF-L-vFc $^{\prime\prime}$ Figure 2B.

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	ctg L	O P O	а В В	tgc C	agc S	ct L	a C C C	gaa E	t c t	gag E	дда G
	aag K -20	a C C H H	gtg V	ctg L	ctg L	990 G	CCC P	atg M	gcc A	ctg L	ggt G
	atg M	gcc -1	саа О 20	аад К 40	ccc P 60	адс 80	ggt G 100	cag Q 120	ttc F	ttc F 160	t c c
	000 P	gaa E	gag E	tac Y	gct A	cat H	ttg L	cag Q	gcc A	agc S	ggt G
	agc S	cag Q	tta L	a C C	tgg W	ctc L	gag E	tgg W	ccg P	cag Q	ggc G
	cag Q	gtg V	t C C	gcc A	O O O	ca Q	OCC P	atc H	atg M	ctg L	ggt G
	a T	а Н а	aag K	tgt C	a T L	agc S	tcc s	a T	gcc A	cat H	tcc S
	gcc	tgg W	ctc L	ctg L	990 G	ttg L	atc I	acc T	ggt G	tcc S	gga G
	cct P	ctc	ctg L	aag K	ctg L	tgc C	999 G	gcc A	cag Q	gcc	CCC P
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SEQ 3	atg M	Cac H H	Cag D	ctc	g G G	ctg L	gcc A	gcc A	cag Q	gtc	ctt L
	O O	tgg W	CCC P	gcg AD	C C C	сад Д	Cag OO	gtc 'C		999 999 6	Cac
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d Seg	U U	ctg L	tcc s	990 G	gtg V	gcc A	ct c	ctg L	cct P	gca A	cta L
DNA Amino Acid Sequence	ctt IIII	ctg L	agc S	gat D	ctg L	cag Q	999 G	cag Q	gcc A	cgg R	gtt V
DNA Amino	aag ctt <i>Hin</i> dIII	cag Q	gcc A	gg c C	gag E	გ ი ი	cag Q	ctg L	atg M	cgc R	aga R

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gca A	ctc L	CCC P	ccg P	cag Q	t c c	ctg L	33c G	tac Y	a GC	gct	
O O O	act	gac D	aag K	CaC H	t S	a G G	aaa K	a a C	ct a L	gag E	a ttc CoRI
tgc C	gac D	gaa E	aca H	ctg L	ccg P	tac Y	gtc V	a a C	agg R	cat H	gaa Ec
G ca 200	aag K 220	cag Q 240	aag K 260	gtc V 280	ctc L 300	gtg V 320	ctg L 340	gag E 360	agc S 380	atg M 400	tga
CC a	CCC P	agc S	gcc A	a T	ggc G	cag Q	tgc C	ccg P	tac Y	gtg V	aaa K 419
tgc C	a A A	gtg V	a N N	ctc L	aaa K	cca P	a T	cag Q	ctc L	t S	ggt G
СС Р	cca P	gac D	cat H	gtc V	a a N	gag E	ctg L	999 6	t F F	tgc C	ctg L
CC P	O P O	gtg V	gtg V	a S S	t S	cga R	agc S	aa N	tt F	t S S	tct S
ggt G	t F	gtg V	gag E	gtc V	gtc V	CCC P	gtc V	a လ လ	tcc s	tt F	ctg L
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tcc S	gtc V	acg T	gat D	tac Y	aag K	a R R	aag K	gag E	t S	ggg G	agc S
gag E	tca S	gtc V	gtg V	acg T	t Y	gcc	acc	gtg V	gac D	gag E	aag K
tca S 190	cca P 210	gag E 230	tac Y 250	agc S 270	gag Е 290	aaa K 310	atg M 330	9cc A 350	ctg L 370	င်ရဋ္ဌ (၃) (၁)	cag 2 410
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DNA and deduced amino acid sequences of hG-CSF-L-vFc $_{\gamma 1}$ Figure 2C.

	DNA Amino Acid Sequence		SEQ II	ID NO:	21													
ttc ctg ttg cc cag agt cc ttg aca gtg cag gaa gcc ccc ctg ggc cct ll l l l l l l l l l l l l l l l l	e CCC			13	დ დ				-	S S	S P	atg å M					ctg L	09
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ccg P	a A M	Ca H	a A M	gtc	ctc	gtg V	ctg L	gag E	agc S	atg M	tga
cca P	0 0 0	agc S	gcc	a G	gcc A	cag Q	tgc C	ccg P	tac Y	gtg V	aaa K 417
tgc C	a N N	gtg V	a N N	ct c	aaa K	CC P	a G	cag Q	ctc	t S S	ggt G
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